#5

OIPE

RAW SEQUENCE LISTING   DATE: 11/14/2001
PATENT APPLICATION: US/09/757,982A   TIME: 15:36:21

Input Set : N:\Crf3\RULE60\09757982A.txt
Output Set: N:\CRF3\11142001\I757982A.raw

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   3 <110> APPLICANT: Acton, Susan
   5 <120> TITLE OF INVENTION: NOVEL PROTEIN KINASE MOLECULES AND USES THEREFOR
   7 <130> FILE REFERENCE: MNI-050
C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/757,982A
  10 <141> CURRENT FILING DATE: 2001-01-10
  12 <150> PRIOR APPLICATION NUMBER: 09/163,115
  13 <151> PRIOR FILING DATE: 1998-09-29
  15 <160> NUMBER OF SEQ ID NOS: 15
  17 <170> SOFTWARE: PatentIn Ver. 2.0
  19 <210> SEQ ID NO: 1
  20 <211> LENGTH: 4137
  21 <212> TYPE: DNA
  22 <213> ORGANISM: Homo sapiens
  24 <220> FEATURE:
  25 <223> OTHER INFORMATION: 'n' at positions 2872, 3597 and 3682 may be any nucleic acid
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  28 <221> NAME/KEY: CDS
  29 <222> LOCATION: (297)..(1202)
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  41                                                                Met
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  52 ttt cga ata gaa aag aaa att ggt cgc gga caa ttt agt gaa gtt tat   443
  53 Phe Arg Ile Glu Lys Lys Ile Gly Arg Gly Gln Phe Ser Glu Val Tyr
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  56 aga gca gcc tgt ctc ttg gat gga gta cca gta gct tta aaa aaa gtg   491
  57 Arg Ala Ala Cys Leu Leu Asp Gly Val Pro Val Ala Leu Lys Lys Val
  58  50                  55                  60                  65
  60 cag ata ttt gat tta atg gat gcc aaa gca cgt gct gat tgc atc aaa   539
  61 Gln Ile Phe Asp Leu Met Asp Ala Lys Ala Arg Ala Asp Cys Ile Lys
  62              70                  75                  80
  64 gaa ata gat ctt ctt aag caa ctc aac cat cca aat gta ata aaa tat   587
  65 Glu Ile Asp Leu Leu Lys Gln Leu Asn His Pro Asn Val Ile Lys Tyr
  66              85                  90                  95
  68 tat gca tca ttc att gaa gat aat gaa cta aac ata gtt ttg gaa cta   635
  69 Tyr Ala Ser Phe Ile Glu Asp Asn Glu Leu Asn Ile Val Leu Glu Leu
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Input Set : N:\Crf3\RULE60\09757982A.txt
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76 aag agg cta att cct gaa aga act gtt tgg aag tat ttt gtt cag ctt   731
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80 tgc agt gca ttg gaa cac atg cat tct cga aga gtc atg cat aga gat   779
81 Cys Ser Ala Leu Glu His Met His Ser Arg Arg Val Met His Arg Asp
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84 ata aaa cca gct aat gtg ttc att aca gcc act ggg gtg gta aaa ctt   827
85 Ile Lys Pro Ala Asn Val Phe Ile Thr Ala Thr Gly Val Val Lys Leu
86             165             170             175
88 gga gat ctt ggg ctt ggc cgg ttt ttc agc tca aaa acc aca gct gca   875
89 Gly Asp Leu Gly Leu Gly Arg Phe Phe Ser Ser Lys Thr Thr Ala Ala
90             180             185             190
92 cat tct tta gtt ggt acg cct tat tac atg tct cca gag aga ata cat   923
93 His Ser Leu Val Gly Thr Pro Tyr Tyr Met Ser Pro Glu Arg Ile His
94     195             200             205
96 gaa aat gga tac aac ttc aaa tct gac atc tgg tct ctt ggc tgt cta   971
97 Glu Asn Gly Tyr Asn Phe Lys Ser Asp Ile Trp Ser Leu Gly Cys Leu
98 210             215             220             225
100 cta tat gag atg gct gca tta caa agt cct ttc tat ggt gac aaa atg   1019
101 Leu Tyr Glu Met Ala Ala Leu Gln Ser Pro Phe Tyr Gly Asp Lys Met
102             230             235             240
104 aat tta tac tca ctg tgt aag aag ata gaa cag tgt gac tac cca cct   1067
105 Asn Leu Tyr Ser Leu Cys Lys Lys Ile Glu Gln Cys Asp Tyr Pro Pro
106             245             250             255
108 ctt cct tca gat cac tat tca gaa gaa ctc cga cag tta gtt aat atg   1115
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112 tgc atc aac cca gat cca gag aag cga cca gac gtc acc tat gtt tat   1163
113 Cys Ile Asn Pro Asp Pro Glu Lys Arg Pro Asp Val Thr Tyr Val Tyr
114     275             280             285
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241                  85                  90                  95
243 Tyr Tyr Ala Ser Phe Ile Glu Asp Asn Glu Leu Asn Ile Val Leu Glu
244             100                 105                 110
246 Leu Ala Asp Ala Gly Asp Leu Ser Arg Met Ile Lys His Phe Lys Lys
247             115                 120                 125
249 Gln Lys Arg Leu Ile Pro Glu Arg Thr Val Trp Lys Tyr Phe Val Gln
250         130                 135                 140
252 Leu Cys Ser Ala Leu Glu His Met His Ser Arg Arg Val Met His Arg
253 145                 150                 155                 160
255 Asp Ile Lys Pro Ala Asn Val Phe Ile Thr Ala Thr Gly Val Val Lys
256                 165                 170                 175
258 Leu Gly Asp Leu Gly Leu Gly Arg Phe Phe Ser Ser Lys Thr Thr Ala
259             180                 185                 190
261 Ala His Ser Leu Val Gly Thr Pro Tyr Tyr Met Ser Pro Glu Arg Ile
262         195                 200                 205
264 His Glu Asn Gly Tyr Asn Phe Lys Ser Asp Ile Trp Ser Leu Gly Cys
265         210                 215                 220
267 Leu Leu Tyr Glu Met Ala Ala Leu Gln Ser Pro Phe Tyr Gly Asp Lys
268 225                 230                 235                 240
270 Met Asn Leu Tyr Ser Leu Cys Lys Lys Ile Glu Gln Cys Asp Tyr Pro
271             245                 250                 255
273 Pro Leu Pro Ser Asp His Tyr Ser Glu Glu Leu Arg Gln Leu Val Asn
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319                 100                 105                 110
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322 Leu Ala Asp Ala Gly Asp Leu Ser Arg Met Ile Lys His Phe Lys Lys
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325 caa aag agg cta att cct gaa aga act gtt tgg aag tat ttt gtt cag   432
326 Gln Lys Arg Leu Ile Pro Glu Arg Thr Val Trp Lys Tyr Phe Val Gln
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331 145                 150                 155                 160
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351 225                 230                 235                 240
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354 Met Asn Leu Tyr Ser Leu Cys Lys Lys Ile Glu Gln Cys Asp Tyr Pro
355                 245                 250                 255
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363         275                 280                 285
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371 <211> LENGTH: 2120
372 <212> TYPE: DNA
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VERIFICATION SUMMARY   DATE: 11/14/2001
PATENT APPLICATION: US/09/757,982A   TIME: 15:36:22

Input Set : N:\Crf3\RULE60\09757982A.txt
Output Set: N:\CRF3\11142001\I757982A.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application Number
L:174 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:198 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:202 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1